

Serial Number: 09/844,864**ENTERED**

CRF Processing Date:

5/11/2001

Edited by:

Verified by:

(STIC staff)

Changed a file from non-ASCII to ASCII

Changed the margins in cases where the sequence text was "wrapped" down to the next line.

Edited a format error in the Current Application Data section, specifically:

Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other _____

Added the mandatory heading and subheadings for "Current Application Data".

Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

Changed the spelling of a mandatory field (the headings or subheadings), specifically:

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

Inserted colons after headings/subheadings. Headings edited included:

Deleted extra, invalid, headings used by an applicant, specifically:

Deleted: non-ASCII "garbage" at the beginning/end of file\$; secretary initials/filename at end of file;
 page numbers throughout text; other invalid text, such as _____

Inserted mandatory headings, specifically: _____

Corrected an obvious error in the response, specifically: _____

Edited identifiers where upper case is used but lower case is required, or vice versa.

Corrected an error in the Number of Sequences field, specifically: _____

A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

Other:

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/844,864

DATE: 05/23/2001
TIME: 13:11:58

Input Set : A:\Pto.amc
Output Set: C:\CRF3\05232001\I844864.raw

2 <110> APPLICANT: Matzuk, Martin
4 Ren, Yongsheng
6 Wu, Xuemei
10 <120> TITLE OF INVENTION: OVARY SPECIFIC GENES AND PROTEINS
14 <130> FILE REFERENCE: P01925US2 / 09807797 / OTA 99-48
C--> 18 <140> CURRENT APPLICATION NUMBER: US/09/844,864
20 <141> CURRENT FILING DATE: 2001-04-27
24 <150> PRIOR APPLICATION NUMBER: 60/106,020
26 <151> PRIOR FILING DATE: 1998-10-28
30 <150> PRIOR APPLICATION NUMBER: PCT/US99/25209
32 <151> PRIOR FILING DATE: 1999-10-28
36 <160> NUMBER OF SEQ ID NOS: 25
40 <170> SOFTWARE: PatentIn version 3.0
44 <210> SEQ ID NO: 1
46 <211> LENGTH: 1277
48 <212> TYPE: DNA
50 <213> ORGANISM: Mus musculus
54 <400> SEQUENCE: 1
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59 cgacccgcgc ccccttcctt cctcccccggc tacagacagc tcacggccgc ggagtacgtc 180
61 gacagccacc agcgggcaca gctcatggcc ctgctgtcgc ggatgggtcc cgggtcggc 240
63 agcagccgtg acgctgcgtt gcaggtgaac ccgcgcggcg acgcctcggt gcagtgttca 300
65 ctcgggcgcc gcacgctgca gcctgcaggg tgccgagcca gccccgacgc ccgatcggtt 360
67 tcctgtcaac cccgtggcca cgccggcgcc gggagatccc cgcgatcctg gcagaccgt 420
69 gccccgttct cgtccgtgac cttctgtggc ctctccttct cactggaggt tgcgggaggc 480
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73 ccgagagagg tggccgcgag gaaagcggtc ccccaagccgc gaagcgagga gggcgatgtt 600
75 caggtgcag ggcaggccgg gtgggagcag cagccaccac cggaggaccg gaacagtgt 660
77 gcggcgatgc agtctgagcc tgggagcag gagccatgtc ctgcccaga gatggctcag 720
79 gaccccggtt attcggatgc ccctcgagac caggcctccc cgaaacgac ggacgaggac 780
81 aaggagcggc tgcgttcca gttcttagag cagaagtacg gctactatca ctgcaaggac 840
83 tgcaaaatcc ggtgggagag cgcttatgtg tgggtgtgc agggcaccag taaggttta 900
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87 cacctgtcaa agttgtaaaa gaactagatg tgcctgccc gtcagattc gccacgtgga 1020
89 ccctaaacgc ccccatcgcc aagacttgtg tgggagatgc aaggacaaac gcctgtcctg 1080
91 cgacagcacc tttagttca aatacatcat tttagtggagatc tcgaaaacgt ttctgtttaga 1140
93 tggggctaat ggaatggaca agtgagctt ctcccctttt cacctcttcc ctttccaaat 1200
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104 <212> TYPE: PRT
106 <213> ORGANISM: Mus musculus
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116 20 25 30
118 Pro Ala Pro Pro Ser Phe Leu Pro Gly Tyr Arg Gln Leu Met Ala Ala
119 35 40 45
121 Glu Tyr Val Asp Ser His Gln Arg Ala Gln Leu Met Ala Leu Leu Ser
122 50 55 60
124 Arg Met Gly Pro Arg Ser Val Ser Ser Arg Asp Ala Ala Val Gln Val
125 65 70 75 80
127 Asn Pro Arg Arg Asp Ala Ser Val Gln Cys Ser Leu Gly Arg Arg Thr
128 85 90 95
130 Leu Gln Pro Ala Gly Cys Arg Ala Ser Pro Asp Ala Arg Ser Gly Ser
131 100 105 110
133 Cys Gln Pro Arg Gly His Ala Gly Ala Gly Arg Ser Pro Arg Ser Trp
134 115 120 125
136 Gln Thr Val Ala Pro Phe Ser Ser Val Thr Phe Cys Gly Leu Ser Ser
137 130 135 140
139 Ser Leu Glu Val Ala Gly Gly Arg Gln Thr Pro Thr Lys Gly Glu Gly
140 145 150 155 160
142 Ser Pro Ala Ser Ser Gly Thr Arg Glu Pro Glu Pro Arg Glu Val Ala
143 165 170 175
145 Ala Arg Lys Ala Val Pro Gln Pro Arg Ser Glu Glu Gly Asp Val Gln
146 180 185 190
148 Ala Ala Gly Gln Ala Gly Trp Glu Gln Gln Pro Pro Pro Glu Asp Arg
149 195 200 205
151 Asn Ser Val Ala Ala Met Gln Ser Glu Pro Gly Ser Glu Glu Pro Cys
152 210 215 220
154 Pro Ala Ala Glu Met Ala Gln Asp Pro Gly Asp Ser Asp Ala Pro Arg
155 225 230 235 240
157 Asp Gln Ala Ser Pro Gln Ser Thr Glu Gln Asp Lys Glu Arg Leu Arg
158 245 250 255
160 Phe Gln Phe Leu Glu Gln Lys Tyr Gly Tyr Tyr His Cys Lys Asp Cys
161 260 265 270
163 Lys Ile Arg Trp Glu Ser Ala Tyr Val Trp Cys Val Gln Gly Thr Ser
164 275 280 285
166 Lys Val Tyr Phe Lys Gln Phe Cys Arg Val Cys Glu Lys Ser Tyr Asn
167 290 295 300
169 Pro Tyr Arg Val Glu Asp Ile Thr Cys Gln Ser Cys Lys Arg Thr Arg
170 305 310 315 320
172 Cys Ala Cys Pro Val Arg Phe Arg His Val Asp Pro Lys Arg Pro His
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183 <211> LENGTH: 1817
185 <212> TYPE: DNA
187 <213> ORGANISM: Mus musculus
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196	cagagtctac	tgagggatga	ggccttggcc	atttctgctc	tcacggacct	gccccagagt	180									
198	ctgttcccag	taattttga	ggaggccttc	actgatggat	atataaggat	cttgaaggcc	240									
200	atgatacctg	tgtggccctt	cccatacctt	tctttaggaa	agcagataaa	taattgcaac	300									
202	ctggagactt	tgaaggctat	gcttgaggga	ctagatatac	tgcttgaca	aaaggttcaa	360									
204	accagtaggt	gcaaactcag	agtaattaaat	tggagagaag	atgacttgaa	gatatgggct	420									
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208	agtgctggct	gtgaggtgaa	gaaagaattg	aaggtgacga	ctgaagtctt	tcgcataaag	540									
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214	atcttcaaaa	ctgtacacgc	agactgtata	caggagctt	tccttaagatg	tatctgcata	720									
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218	ctagatcaca	tcataggtac	cttcagttt	ggtgattctg	aaaagcttga	tgaggagaca	840									
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236	tctagagtgc	tatgatgaca	gtgggttaat	actaacacac	agattagaaa	gtttttgtcc	1380									
238	ttagcttctg	gatatactga	gagccaaaag	acagctccat	agtgtctcct	ttcaaacaac	1440									
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271					20			25			30					
273	Gln	Ser	Leu	Leu	Arg	Asp	Glu	Ala	Leu	Ala	Ile	Ser	Ala	Leu	Thr	Asp
274					35			40			45					
276	Leu	Pro	Gln	Ser	Leu	Phe	Pro	Val	Ile	Phe	Glu	Glu	Ala	Phe	Thr	Asp
277					50			55			60					
279	Gly	Tyr	Ile	Gly	Ile	Leu	Lys	Ala	Met	Ile	Pro	Val	Trp	Pro	Phe	Pro
280	65					70			75			80				
282	Tyr	Leu	Ser	Leu	Gly	Lys	Gln	Ile	Asn	Asn	Cys	Asn	Leu	Glu	Thr	Leu
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285	Lys	Ala	Met	Leu	Glu	Gly	Leu	Asp	Ile	Leu	Leu	Ala	Gln	Lys	Val	Gln

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291	Lys Ile Trp Ala Gly Ser His Glu Gly Glu Gly Leu Pro Asp Phe Arg			
292	130	135	140	
294	Thr Glu Lys Gln Pro Ile Glu Asn Ser Ala Gly Cys Glu Val Lys Lys			
295	145	150	155	160
297	Glu Leu Lys Val Thr Thr Glu Val Leu Arg Met Lys Gly Arg Leu Asp			
298	165	170	175	
300	Glu Ser Thr Thr Tyr Leu Leu Gln Trp Ala Gln Gln Arg Lys Asp Ser			
301	180	185	190	
303	Ile His Leu Phe Cys Arg Lys Leu Leu Ile Glu Gly Leu Thr Lys Ala			
304	195	200	205	
306	Ser Val Ile Glu Ile Phe Lys Thr Val His Ala Asp Cys Ile Gln Glu			
307	210	215	220	
309	Leu Ile Leu Arg Cys Ile Cys Ile Glu Glu Leu Ala Phe Leu Asn Pro			
310	225	230	235	240
312	Tyr Leu Lys Leu Met Lys Ser Leu Phe Thr Leu Thr Leu Asp His Ile			
313	245	250	255	
315	Ile Gly Thr Phe Ser Leu Gly Asp Ser Glu Lys Leu Asp Glu Glu Thr			
316	260	265	270	
318	Ile Phe Ser Leu Ile Ser Gln Leu Pro Thr Leu His Cys Leu Gln Lys			
319	275	280	285	
321	Leu Tyr Val Asn Asp Val Pro Phe Ile Lys Gly Asn Leu Lys Glu Tyr			
322	290	295	300	
324	Leu Arg Cys Leu Lys Lys Pro Leu Glu Thr Leu Cys Ile Ser Asn Cys			
325	305	310	315	320
327	Asp Leu Ser Gln Ser Asp Leu Asp Cys Leu Pro Tyr Cys Leu Asn Ile			
328	325	330	335	
330	Cys Glu Leu Lys His Leu His Ile Ser Asp Ile Tyr Leu Cys Asp Leu			
331	340	345	350	
333	Leu Leu Glu Pro Leu Gly Phe Leu Leu Glu Arg Val Gly Asp Thr Leu			
334	355	360	365	
336	Lys Thr Leu Glu Leu Asp Ser Cys Cys Ile Val Asp Phe Gln Phe Ser			
337	370	375	380	
339	Ala Leu Leu Pro Ala Leu Ser Gln Cys Ser His Leu Arg Glu Val Thr			
340	385	390	395	400
342	Phe Tyr Asp Asn Asp Val Ser Leu Pro Phe Leu Lys Thr Thr Ser Thr			
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346	420	425		
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350	<211> LENGTH: 1018			
352	<212> TYPE: DNA			
354	<213> ORGANISM: Mus musculus			
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Input Set : A:\Pto.amc
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 369 tctgcctggg ggagaaagcc aaagaggagg tgaaccgtgt ggaagtcctc tcccaggaag 360
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 381 aaaagaaggt ggaaaaagaa gaggatgaaa cagtagtgag gcccagccct caggacaaga 720
 383 gtccctggaa gaaggagaaa tctacaccca gagcaaagaa gccagtgacc aagaaatgac 780
 385 ctcatcttag catcttctgc gtccaaggca ggatgtccag cagctgtgtt ttgggtgcagg 840
 387 tgtccagccc caccacccta gtctgaatgt aataagggtgg tgtggctgta accctgtaac 900
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 412 Arg Gly Gln Gly Glu Lys Lys Asp Ser Cys Lys Leu Leu Ser Thr
 413 35 40 45
 415 Ile Cys Leu Gly Glu Lys Ala Lys Glu Glu Val Asn Arg Val Glu Val
 416 50 55 60
 418 Leu Ser Gln Glu Gly Arg Lys Pro Pro Ile Thr Ile Ala Thr Leu Lys
 419 65 70 75 80
 421 Ala Ser Val Leu Pro Met Val Thr Val Ser Gly Ile Glu Leu Ser Pro
 422 85 90 95
 424 Pro Val Thr Phe Arg Leu Arg Thr Gly Ser Gly Pro Val Phe Leu Ser
 425 100 105 110
 427 Gly Leu Glu Cys Tyr Glu Thr Ser Asp Leu Thr Trp Glu Asp Asp Glu
 428 115 120 125
 430 Glu Glu Glu Glu Glu Glu Glu Glu Asp Glu Asp Glu Asp Ala Asp
 431 130 135 140
 433 Ile Ser Leu Glu Glu Ile Pro Val Lys Gln Val Lys Arg Val Ala Pro
 434 145 150 155 160
 436 Gln Lys Gln Met Ser Ile Ala Lys Lys Lys Val Glu Lys Glu Glu
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 445 <210> SEQ ID NO: 7
 447 <211> LENGTH: 214
 449 <212> TYPE: DNA
 451 <213> ORGANISM: Mus musculus

VERIFICATION SUMMARY
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L:18 M:270 C: Current Application Number differs, Replaced Current Application Number